## Controlling microbial co-culture populations based on substrate pulsing can lead to stability through differential fitness advantages.

 $J.Andres Martinez^1, Matheo Delvenne^{1,Y}, Lucas Henrion^{1,Y}, Fabian Moreno^{1,Y}, Samuel Telek^1, Christian Dusny^2, Frank Delvigne^{1,*}$ 

**1** TERRA Research and Teaching Centre, Microbial Processes and Interactions (MiPI), Gembloux Agro-Bio Tech, University of Liége, Gembloux, Belgium.

**2** Microbial Single Cell Analysis, Department of Solar Materials, Helmholtz-Centre for Environmental Research-UFZ Leipzig, Permoserstr. 15 04318 Leipzig, Germany.

\* f.delvigne@uliege.be

## Supplementary File 4: Continuous and Discontinuous culture supplementary figures

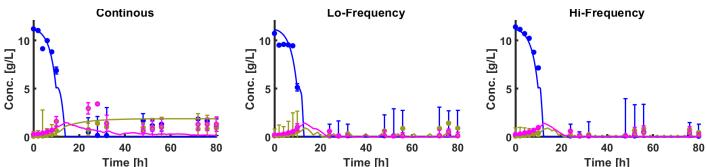


Fig 1. Measurement data set coverage for the Continuous, Low-frequency and High-frequency feed co-culture experiments from time 24 to 80.

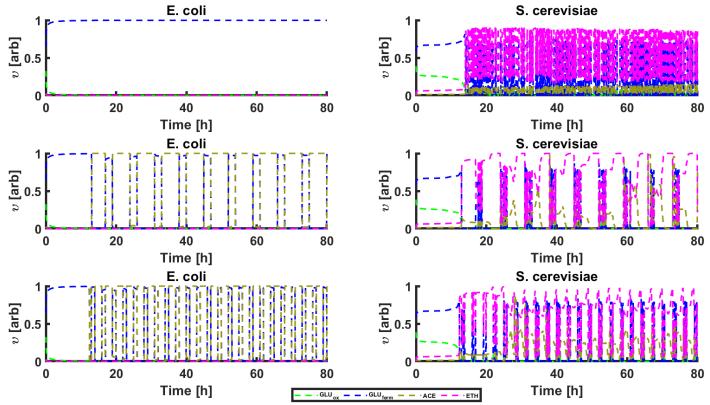


Fig 2. Cybernetic variable v for E. coli and S. cerevisiae during the simulations made for Continuous culture (up), low frequency (middle) and high frequency (down) pulsing experiments.

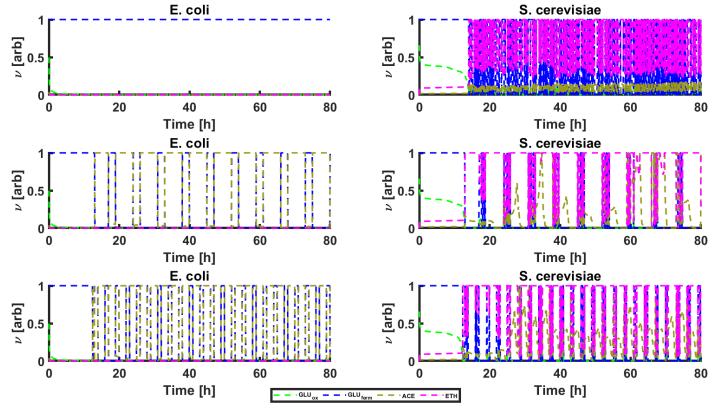


Fig 3. Cybernetic variable  $\nu$  for *E. coli* and *S. cerevisiae* during the simulations made for Continuous culture (up), low frequency (middle) and high frequency (down) pulsing experiments.

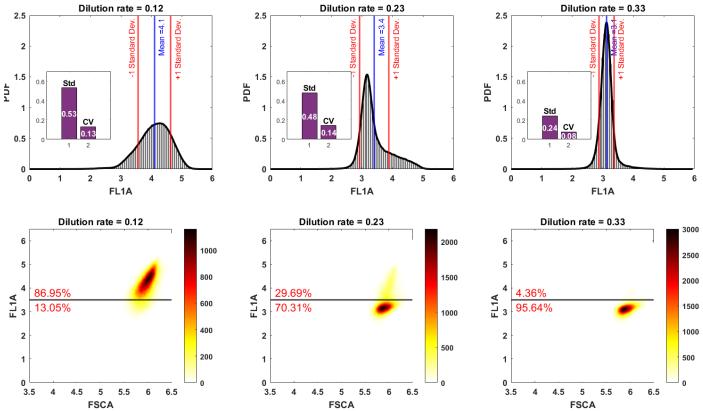


Fig 4. top row: FSCA Probability density function, mean, standard deviations and coefficient of variation for the axenic continuous cultures of *S. cerevisiae* at dilution rates of 0.12, 0.23 and 0.33  $h^{-1}$ , respectively. bottom row: Flow citometry data for the triplicates at the different dilution rates and their  $GFP^+$  and  $GFP^-$  percentual calculations along the threshold at 3.5.

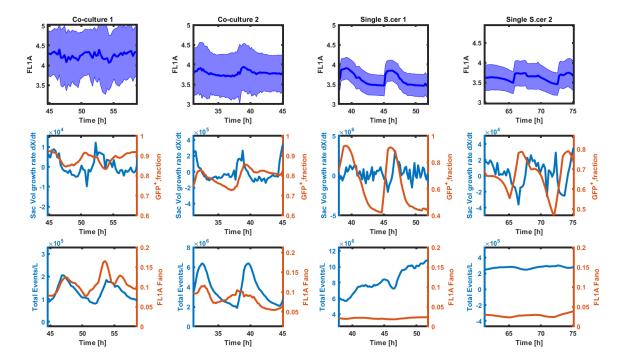


Fig 5. Time evolution of FL1, approximated growth rate,  $GFP^+$  fraction, total events over liter and Fano factor across example cycles for the low-frequency feed regime experiments. Plotted values correspond for *S. cerevisiae* population in coculture and in single culture (duplicates)